DOCUMENT TYPE: Patent English LANGUAGE:

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.		DATE
US 2005003369	A1	20050106	US 2003-684205		20031010
US 2005123987	A1	20050609	US 2005-40759		20050121
PRIORITY APPLN. INFO.:			US 2002-417803P	P	20021010
			US 2002-417817P	P	20021011
			US 2003-684205	A1	20031010

9050-76-4, RNase H IT

> RL: ARU (Analytical role, unclassified); BSU (Biological study, unclassified); ANST (Analytical study); BIOL (Biological study) (method for depleting specific nucleic acids from mixture isolated from blood)

L25 ANSWER 8 OF 236 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 1

ACCESSION NUMBER: 2005:1243337 CAPLUS

Specific recognition and cleavage of the plus-strand TITLE:

primer by reverse transcriptase

Atwood-Moore, Angela; Ejebe, Kenechi; Levin, Henry L. AUTHOR (S):

CORPORATE SOURCE: Section on Eukaryotic Transposable Elements, Laboratory of Gene Regulation and Development, National Institute of Child Health and Human

Development, National Institutes of Health, Bethesda,

MD, 20892, USA

SOURCE: Journal of Virology (2005), 79(23), 14863-14875

CODEN: JOVIAM; ISSN: 0022-538X American Society for Microbiology

DOCUMENT TYPE: Journal LANGUAGE: English

PUBLISHER:

REFERENCE COUNT: THERE ARE 35 CITED REFERENCES AVAILABLE FOR THIS 35

RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

Reverse transcriptases (RTs) of retroviruses and long terminal repeat AB (LTR) -retrotransposons possess DNA polymerase and RNase H activities. During reverse transcription these activities are necessary for the programmed sequence of events that include template switching and primer processing. Integrase then inserts the completed cDNA into the genome of the host cell. The RT of the LTR-retrotransposon Tf1 was subjected to random mutagenesis, and the resulting transposons were screened with genetic assays to test which mutations reduced reverse transcription and which inhibited integration. We identified a cluster of mutations in the RNase H domain of RT that were surprising because they blocked integration without reducing cDNA levels. The results of immunoblots demonstrated that these mutations did not reduce levels of RT or integrase. DNA blots showed that the mutations did not lower the amts. of full-length cDNA. The sequences of the 3' ends of the cDNA revealed that mutations within the cluster in RNase H specifically reduced the removal of the polypurine tract (PPT) primer from the ends of the These results indicate that primer removal is not a necessary component of reverse transcription. The residues mutated in Tf1 RNase H are conserved in human immunodeficiency virus type 1 and make direct contact with DNA opposite the PPT. Thus, our results identify a conserved element in RT that contacts the PPT and is specifically required for PPT removal.

=> d hist full

(FILE 'HOME' ENTERED AT 16:26:43 ON 09 JAN 2006)

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FILE 'HOME' ENTERED AT 16:26:56 ON 09 JAN 2006 SET LINE LOGIN SET DETAIL LOGIN

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FILE 'CAPLUS, MEDLINE, BIOSIS' ENTERED AT 16:27:10 ON 09 JAN 2006
            34 SEA ABB=ON PLU=ON BAIT (2A) (MOLECULE OR PROBE OR PRIMER)
L1
             O SEA ABB=ON PLU=ON L1 AND GLOBIN
L2
            118 SEA ABB=ON PLU=ON BAIT (10A) (MOLECULE OR PROBE OR PRIMER OR
L3
                OLIGONUCLEOTIDE?)
              O SEA ABB=ON PLU=ON L3 AND GLOBIN
L4
         81147 SEA ABB=ON PLU=ON COMPLEX? (10A) (REDUC?)
L5
           132 SEA ABB=ON PLU=ON L5 AND GLOBIN
L6
             O SEA ABB=ON PLU=ON L3 AND HOUSEKEEPING
L7
              6 SEA ABB=ON PLU=ON L3 AND ABUNDAN?
L8
          7946 SEA ABB=ON PLU=ON GLOBIN AND (REDUC? OR DEPLET? OR REMOV?)
L9
          2542 SEA ABB=ON PLU=ON GLOBIN(S) (REDUC? OR DEPLET? OR REMOV?)
L10
L11
          2504 SEA ABB=ON PLU=ON GLOBIN(30A) (REDUC? OR DEPLET? OR REMOV?)
           674 SEA ABB=ON PLU=ON GLOBIN(30A)(REDUC? OR DEPLET? OR REMOV?)(S)
L12
                (MRNA OR EXPRESSION OR RNA)
           674 SEA ABB=ON PLU=ON GLOBIN(30A) (REDUC? OR DEPLET? OR REMOV?) (30
L13
                A) (MRNA OR EXPRESSION OR RNA)
L14
            310 DUP REM L13 (364 DUPLICATES REMOVED)
L15
            31 SEA ABB=ON PLU=ON L14 AND (RNASE OR DIGEST?)
L16
            31 DUP REM L15 (0 DUPLICATES REMOVED)
L17
            23 DUP REM L1 (11 DUPLICATES REMOVED)
L18
            37 SEA ABB=ON PLU=ON L3 AND (MRNA OR CDNA OR RNA)
L19
            25 DUP REM L18 (12 DUPLICATES REMOVED)
L20
             7 SEA ABB=ON PLU=ON RNA(10A) STABILIZER? AND (BLOOD OR PLASMA
               OR SERUM)
             5 DUP REM L20 (2 DUPLICATES REMOVED)
L21
L22
             O SEA ABB=ON PLU=ON L17 AND BLOOD
L23
             4 SEA ABB=ON PLU=ON L18 AND BLOOD
L24
           500 SEA ABB=ON PLU=ON (DEPLET? OR REMOV? OR INACTIV?) (20A) (RNASE(
                2A) H)
           236 DUP REM L24 (264 DUPLICATES REMOVED)
L25
            22 SEA ABB=ON PLU=ON L25 AND (HEAT? OR COLUMN? OR EDTA)
L26
L27
             22 DUP REM L26 (0 DUPLICATES REMOVED)
               D L16 TI 1-10
               D L16 TI 11-20
               D L16 TI 21-31
               D L17 TI 1-10
               D L17 TI 11-23
               D L25 TI 1-12
               D L25 TI 13-23
               D L19 TI 1-12
               D L19 TI 13-25
               D L21 TI 1-10
               D L16 IBIB KWIC 1,3,4
               D L16 IBIB KWIC 12-14
               D L16 IBIB KWIC 23,28,30
               D L17 IBIB KWIC 7
               D L19 IBIB KWIC 3,13
               D L25 IBIB KWIC 3,4,8
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FILE HOME

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FILE MEDLINE

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On December 11, 2005, the 2006 MeSH terms were loaded.

The MEDLINE reload for 2006 will soon be available. For details on the 2005 reload, enter HELP RLOAD at an arrow promt (=>). See also:

http://www.nlm.nih.gov/mesh/

http://www.nlm.nih.gov/pubs/techbull/nd04/nd04_mesh.html

http://www.nlm.nih.gov/pubs/techbull/nd05/nd05_med_data_changes.html

http://www.nlm.nih.gov/pubs/techbull/nd05/nd05_2006_MeSH.html

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MEDLINE thesauri in the /CN, /CT, and /MN fields incorporate the MeSH 2006 vocabulary.

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FILE COVERS 1969 TO DATE.

CAS REGISTRY NUMBERS AND CHEMICAL NAMES (CNs) PRESENT FROM JANUARY 1969 TO DATE.

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